

In the Claims:

Please amend claim 1 as follows:

1. (amended) A transcription factor with a DNA-binding domain comprising a basic domain and an adjacent Leucine zipper domain with the following amino acids:

L e1 x g1 x x x L e2 x g2 x x x' x L e3 x g3 x x x L e4 x g4 x x x
L e5 (SEQ ID NO: 1)

wherein L is leucine, gi and ei are possible substitution locations, x are any amino acids, and x' is one of tyrosine or glycine and wherein one of the following sets of substituents are at the nine substitution locations ei and gi:

KK KQ AT RQ K	SEQ ID NO: 21
EE EQ AT RQ K	SEQ ID NO: 22
EK KQ AT EE K	SEQ ID NO: 23
EE EQ AT EE K	SEQ ID NO: 24
KK EQ AT RQ K	SEQ ID NO: 25
KE EQ AT RQ K	SEQ ID NO: 26
KK KQ AT EE K	SEQ ID NO: 27
KE EQ AT EE K	SEQ ID NO: 28
EK KQ AT RQ E	SEQ ID NO: 29
KK KQ AT RQ E	SEQ ID NO: 30
EK EQ AT RQ E	SEQ ID NO: 31
EE EQ AT RQ E	SEQ ID NO: 32
EK KQ AT EE E	SEQ ID NO: 33
EE EQ AT EE E	SEQ ID NO: 34
KK SL QE RE K	SEQ ID NO: 35
EK SR QV RE K	SEQ ID NO: 36
KK SR QV RE K	SEQ ID NO: 37,

valine being replaceable by alanine, leucine, methionine, isoleucine, glutamine.